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94

50

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15 Trp Glu Glu Leu Lys Ala Arg Asp His Leu Ala Val Leu Gln Ala Arg  
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Leu Thr Val Ser Gly Leu Lys Ala Arg Ala Lys Lys Asp Glu Leu Gln  
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20 Ile Lys Val Arg Val Asp Met Val Glu Leu Lys Arg Lys Met Met Tyr  
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30 Lys Cys Arg Cys Pro Ser Gln Glu Pro Met Ser Ile Tyr Val Tyr Phe  
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Glu Gln Pro Thr Glu Thr Pro Pro Tyr Ala Tyr Ile Tyr Thr Thr Tyr  
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35 Leu Gln Glu Phe Phe Trp Asp Ala Asn Asp Ile Tyr Tyr Glu Gln His  
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40 Lys Ile Thr Ser Tyr Gln Glu Pro Met Ser Ile Tyr Val Tyr Ser Glu  
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Arg Asn His Glu Val Tyr Glu Gln His Lys Ile Thr Ser Tyr Gln Ala  
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... Val Glu Leu Ile Met Arg Glu Phe Asn Ser Tyr Lys  
115 120 125

Val Ala Ala Leu Phe Lys Ala Arg Asp His Leu Ala Val Val Tyr Val

45

Tyr Leu Gln Ala Arg Leu Thr Val Ser Gly Ile Lys Asp Met Glu

Lys Asp Glu Leu Arg Arg Arg His Arg Gln Asp Ala Leu Ala Arg Val  
210 215 220

5 Tyr Glu Ile Lys Cys Arg Gln Glu Phe Phe Trp Asp Ala Asn Asp Ile  
225 230 235 240

10 Tyr

15 <210> 311

<211> 250

<212> PRT

20 <213> synthetic polyepitope peptide

25 <400> 311

Tyr Val Lys Tyr Val Tyr Glu Ser Phe Asn Leu Val Pro Met Val Ala  
1 5 10 15

30 Thr Val Gln Ile Lys Val Arg Val Asp Met Val Val Leu Glu Glu Thr  
20 25 30

35 Ser Val Met Leu Lys Ala Arg Asp His Leu Ala Val Leu Ala Tyr Ala  
35 40 45

40 Gln Lys Ile Phe Lys Ile Leu Val Thr Glu His Asp Thr Leu Leu Tyr  
50 55 60

Tyr Ala Tyr Ile Tyr Thr Thr Tyr Leu Gln Glu Phe Phe Trp Asp Ala  
65 70 75 80

45 Asn Asp Ile Tyr Gln Glu Pro Met Ser Ile Tyr Val Tyr Arg Ile Phe  
85 90 95

50 Ala Glu Leu Glu Gly Val Ile Ile Tyr Thr Arg Asn His Glu Val Ile  
100 105 110

55 Met Arg Glu Phe Asn Ser Tyr Lys Ser Glu His Pro Thr Phe Thr Ser  
115 120 125

Gln Tyr Gln Ala Arg Leu Thr Val Ser Gly Leu Gln Tyr Asp Pro Val  
130 135 140

5 Ala Ala Leu Phe Tyr Glu Gln His Lys Ile Thr Ser Tyr Gln Ala Ile  
145 150 155 160

10 Arg Glu Thr Val Glu Leu Tyr Val Lys Tyr Val Tyr Glu Ser Phe Asp  
165 170 175

15 Ile Tyr Arg Ile Phe Ala Glu Leu Asn Val Arg Arg Ser Trp Glu Glu  
180 185 190

Leu Lys Ala Arg Ala Lys Lys Asp Glu Leu Glu Leu Lys Arg Lys Met  
195 200 205

20 Met Tyr Met Arg Arg Arg His Arg Gln Asp Ala Leu Ala Arg Val Tyr  
210 215 220

25 Glu Ile Lys Cys Arg Cys Pro Ser Gln Glu Pro Met Ser Ile Tyr Val  
225 230 235 240

30 Tyr Phe Glu Gln Pro Thr Glu Thr Pro Pro  
245 250

<210> 312

35 <211> 241

<212> PRT

40 <213> synthetic polyepitope peptide

<400> 312

45 Asn Val Arg Arg Ser Trp Glu Glu Leu Lys Ala Arg Asp His Leu Ala  
1 5 10 15

50 Val Leu Gln Ala Arg Leu Thr Val Ser Gly Leu Lys Ala Arg Ala Lys  
20 25 30

55 Lys Asp Glu Leu Gln Ile Lys Val Arg Val Asp Met Val Glu Leu Lys  
35 40 45

Arg Lys Met Met Tyr Met Arg Arg His Arg Gln Asp Ala Leu Ala

50

55

60

5 Arg Val Tyr Glu Ile Lys Cys Arg Cys Pro Ser Gln Glu Pro Met Ser  
65 70 75 80

10 Ile Tyr Val Tyr Phe Glu Gln Pro Thr Glu Thr Pro Pro Tyr Ala Tyr  
85 90 95

15 Ile Tyr Thr Thr Tyr Leu Gln Glu Phe Phe Trp Asp Ala Asn Asp Ile  
100 105 110

Tyr Tyr Glu Gln His Lys Ile Thr Ser Tyr Gln Glu Pro Met Ser Ile  
115 120 125

20 Tyr Val Tyr Ser Glu His Pro Thr Phe Thr Ser Gln Tyr Gln Ala Ile  
130 135 140

25 Arg Glu Thr Val Glu Leu Val Thr Glu His Asp Thr Leu Leu Tyr Asn  
145 150 155 160

30 Leu Val Pro Met Val Ala Thr Val Val Leu Glu Glu Thr Ser Val Met  
165 170 175

Leu Arg Ile Phe Ala Glu Leu Glu Gly Val Ile Ile Tyr Thr Arg Asn  
180 185 190

35 His Glu Val Ile Met Arg Glu Phe Asn Ser Tyr Lys Ala Tyr Ala Gln  
195 200 205

40 Lys Ile Phe Lys Ile Leu Gln Tyr Asp Pro Val Ala Ala Leu Phe Tyr  
210 215 220

45 Val Lys Tyr Val Tyr Glu Ser Phe Asp Ile Tyr Arg Ile Phe Ala Glu  
225 230 235 240

Leu

50

<210> 313

55 <211> 241  
<212> PRT

99

## &lt;213&gt; synthetic polyepitope peptide

5 &lt;400&gt; 313

Cys Pro Ser Gln Glu Pro Met Ser Ile Tyr Val Tyr Phe Glu Gln Pro  
1 5 10 15

10

Thr Glu Thr Pro Pro Tyr Ala Tyr Ile Tyr Thr Thr Tyr Leu Gln Glu  
20 25 30

15

Phe Phe Trp Asp Ala Asn Asp Ile Tyr Tyr Glu Gln His Lys Ile Thr  
35 40 45

20

Ser Tyr Gln Glu Pro Met Ser Ile Tyr Val Tyr Ser Glu His Pro Thr  
50 55 60

25

Phe Thr Ser Gln Tyr Val Thr Glu His Asp Thr Leu Leu Tyr Asn Leu  
65 70 75 80

30

Val Pro Met Val Ala Thr Val Val Leu Glu Glu Thr Ser Val Met Leu  
85 90 95

35

Arg Ile Phe Ala Glu Leu Glu Gly Val Ile Ile Tyr Thr Arg Asn His  
100 105 110

40

45

Glu Val Ile Met Arg Glu Phe Asn Ser Tyr Lys Asn Val Arg Arg Ser  
115 120 125

50

55

Trp Glu Glu Leu Lys Ala Arg Asp His Leu Ala Val Leu Gln Ala Arg  
130 135 140

Leu Thr Val Ser Gly Leu Lys Ala Arg Ala Lys Lys Asp Glu Leu Gln  
145 150 155 160

Ile Lys Val Arg Val Asp Met Val Glu Leu Lys Arg Lys Met Met Tyr  
165 170 175

Met Arg Arg Arg His Arg Gln Asp Ala Leu Ala Arg Val Tyr Glu Ile  
180 185 190

Lys Cys Arg Ala Tyr Ala Gln Lys Ile Phe Lys Ile Leu Gln Tyr Asp  
195 200 205

100

Pro Val Ala Ala Leu Phe Tyr Val Lys Tyr Val Tyr Glu Ser Phe Asp  
210 215 220

5 Ile Tyr Arg Ile Phe Ala Glu Leu Gln Ala Ile Arg Glu Thr Val Glu  
225 230 235 240

10 Leu

15 <210> 314

<211> 241

<212> PRT

20 <213> synthetic polyepitope peptide

25 <400> 314

Ala Tyr Ala Gln Lys Ile Phe Lys Ile Leu Asn Leu Val Pro Met Val  
1 5 10 15

30 Ala Thr Val Asn Val Arg Arg Ser Trp Glu Glu Leu Glu Leu Lys Arg  
20 25 30

35 Lys Met Met Tyr Met Val Thr Glu His Asp Thr Leu Leu Tyr Phe Glu  
35 40 45

40 Gln Pro Thr Glu Thr Pro Pro Ala Arg Val Tyr Glu Ile Lys Cys Arg  
50 55 60

Arg Ile Phe Ala Glu Leu Glu Gly Val Val Leu Glu Glu Thr Ser Val  
65 70 75 80

45 Met Leu Cys Pro Ser Gln Glu Pro Met Ser Ile Tyr Val Tyr Ile Ile  
85 90 95

50 Tyr Thr Arg Asn His Glu Val Arg Arg Arg His Arg Gln Asp Ala Leu  
100 105 110

55 Ile Met Arg Glu Phe Asn Ser Tyr Lys Gln Ile Lys Val Arg Val Asp  
115 120 125

101

Met Val Tyr Glu Gln His Lys Ile Thr Ser Tyr Gln Tyr Asp Pro Val  
130 135 140

5 Ala Ala Leu Phe Ser Glu His Pro Thr Phe Thr Ser Gln Tyr Gln Glu  
145 150 155 160

10 Phe Phe Trp Asp Ala Asn Asp Ile Tyr Tyr Val Lys Tyr Val Tyr Glu  
165 170 175

Ser Phe Asp Ile Tyr Arg Ile Phe Ala Glu Leu Lys Ala Arg Asp His  
180 185 190

15

Leu Ala Val Leu Gln Ala Arg Leu Thr Val Ser Gly Leu Lys Ala Arg  
195 200 205

20

Ala Lys Lys Asp Glu Leu Tyr Ala Tyr Ile Tyr Thr Thr Tyr Leu Gln  
210 215 220

25 Glu Pro Met Ser Ile Tyr Val Tyr Gln Ala Ile Arg Glu Thr Val Glu  
225 230 235 240

30

Leu

35 <210> 315

40 <211> 241

<212> PRT

45 <213> synthetic polyepitope peptide

40

50 <400> 315

45 Tyr Glu Gln His Lys Ile Thr Ser Tyr Gln Glu Pro Met Ser Ile Tyr  
1 5 10 15

55 Val Tyr Glu Leu Lys Arg Lys Met Met Tyr Met Arg Arg Arg His Arg  
20 25 30

Gln Asp Ala Leu Ala Arg Val Tyr Glu Ile Lys Cys Arg Ala Tyr Ala  
35 40 45

55 Gln Lys Ile Phe Lys Ile Leu Gln Tyr Asp Pro Val Ala Ala Leu Phe

102

50

55

60

5 Tyr Val Lys Tyr Val Tyr Glu Ser Phe Asp Ile Tyr Arg Ile Phe Ala  
65 70 75 80

10 Glu Leu Ser Glu His Pro Thr Phe Thr Ser Gln Tyr Gln Ala Ile Arg  
85 90 95

10

Glu Thr Val Glu Leu Asn Val Arg Arg Ser Trp Glu Glu Leu Lys Ala  
100 105 110

15

Arg Ala Lys Lys Asp Glu Leu Val Thr Glu His Asp Thr Leu Leu Tyr  
115 120 125

20

Asn Leu Val Pro Met Val Ala Thr Val Val Leu Glu Glu Thr Ser Val  
130 135 140

25

Met Leu Arg Ile Phe Ala Glu Leu Glu Gly Val Ile Ile Tyr Thr Arg  
145 150 155 160

30

Asn His Glu Val Ile Met Arg Glu Phe Asn Ser Tyr Lys Gln Ile Lys  
165 170 175

35

Val Arg Val Asp Met Val Phe Glu Gln Pro Thr Glu Thr Pro Pro Tyr  
180 185 190

40

Ala Tyr Ile Tyr Thr Thr Tyr Leu Gln Glu Phe Phe Trp Asp Ala Asn  
195 200 205

45

Asp Ile Tyr Lys Ala Arg Asp His Leu Ala Val Leu Gln Ala Arg Leu  
210 215 220

50

Thr Val Ser Gly Leu Cys Pro Ser Gln Glu Pro Met Ser Ile Tyr Val  
225 230 235 240

Tyr

55

<210> 316

<211> 241

55

<212> PRT

103

&lt;213&gt; synthetic polyepitope peptide

5    <400> 316

10                Gln Ile Lys Val Arg Val Asp Met Val Ala Arg Val Tyr Glu Ile Lys  
                    1                  5                  10                  15

15                Cys Arg Asn Leu Val Pro Met Val Ala Thr Val Tyr Val Lys Tyr Val  
                    20                  25                  30

20                Tyr Glu Ser Phe Arg Ile Phe Ala Glu Leu Glu Gly Val Tyr Ala Tyr  
                    35                  40                  45

25                Ile Tyr Thr Thr Tyr Leu Ile Ile Tyr Thr Arg Asn His Glu Val Ile  
                    50                  55                  60

30                Met Arg Glu Phe Asn Ser Tyr Lys Ser Glu His Pro Thr Phe Thr Ser  
                    65                  70                  75                  80

35                Gln Tyr Ala Tyr Ala Gln Lys Ile Phe Lys Ile Leu Gln Tyr Asp Pro  
                    85                  90                  95

40                Val Ala Ala Leu Phe Asp Ile Tyr Arg Ile Phe Ala Glu Leu Gln Glu  
                    100                  105                  110

45                Phe Phe Trp Asp Ala Asn Asp Ile Tyr Asn Val Arg Arg Ser Trp Glu  
                    115                  120                  125

50                Glu Leu Lys Ala Arg Asp His Leu Ala Val Leu Val Leu Glu Glu Thr  
                    130                  135                  140

55                Ser Val Met Leu Gln Ala Arg Leu Thr Val Ser Gly Leu Lys Ala Arg  
                    145                  150                  155                  160

60                Ala Lys Lys Asp Glu Leu Glu Leu Lys Arg Lys Met Met Tyr Met Val  
                    165                  170                  175

65                Thr Glu His Asp Thr Leu Leu Tyr Gln Glu Pro Met Ser Ile Tyr Val  
                    180                  185                  190

70                Tyr Arg Arg Arg His Arg Gln Asp Ala Leu Cys Pro Ser Gln Glu Pro  
                    195                  200                  205

104

Met Ser Ile Tyr Val Tyr Phe Glu Gln Pro Thr Glu Thr Pro Pro Tyr  
210 215 220

5 Glu Gln His Lys Ile Thr Ser Tyr Gln Ala Ile Arg Glu Thr Val Glu  
225 230 235 240

10 Leu

15 <210> 317

<211> 241

<212> PRT

20 <213> synthetic polyepitope peptide

25 <400> 317

Asp Ile Tyr Arg Ile Phe Ala Glu Leu Cys Pro Ser Gln Glu Pro Met  
1 5 10 15

30 Ser Ile Tyr Val Tyr Gln Ala Ile Arg Glu Thr Val Glu Leu Lys Ala  
20 25 30

35 Arg Asp His Leu Ala Val Leu Asn Leu Val Pro Met Val Ala Thr Val  
35 40 45

40 Tyr Val Lys Tyr Val Tyr Glu Ser Phe Val Leu Glu Glu Thr Ser Val  
50 55 60

Met Leu Asn Val Arg Arg Ser Trp Glu Glu Leu Ile Met Arg Glu Phe  
65 70 75 80

45 Asn Ser Tyr Lys Ala Tyr Ala Gln Lys Ile Phe Lys Ile Leu Glu Leu  
85 90 95

50 Lys Arg Lys Met Met Tyr Met Gln Tyr Asp Pro Val Ala Ala Leu Phe  
100 105 110

55 Ser Glu His Pro Thr Phe Thr Ser Gln Tyr Gln Ala Arg Leu Thr Val  
115 120 125

105

Ser Gly Leu Lys Ala Arg Ala Lys Lys Asp Glu Leu Gln Ile Lys Val  
130 135 140

5 Arg Val Asp Met Val Arg Arg Arg His Arg Gln Asp Ala Leu Ala Arg  
145 150 155 160

10 Val Tyr Glu Ile Lys Cys Arg Phe Glu Gln Pro Thr Glu Thr Pro Pro  
165 170 175

15 Tyr Ala Tyr Ile Tyr Thr Thr Tyr Leu Gln Glu Phe Phe Trp Asp Ala  
180 185 190

20 Asn Asp Ile Tyr Tyr Glu Gln His Lys Ile Thr Ser Tyr Val Thr Glu  
195 200 205

25 His Asp Thr Leu Leu Tyr Gln Glu Pro Met Ser Ile Tyr Val Tyr Arg  
210 215 220

30 Val

35 <210> 318

40 <211> 241

<212> PRT

<213> synthetic polyepitope peptide

45 Asn Leu Val Pro Met Val Ala Thr Val Gln Ala Arg Leu Thr Val Ser  
1 5 10 15

50 Gly Leu Val Thr Glu His Asp Thr Leu Leu Tyr Tyr Ala Tyr Ile Tyr  
20 25 30

55 Thr Thr Tyr Leu Gln Glu Phe Phe Trp Asp Ala Asn Asp Ile Tyr Val  
35 40 45

Leu Glu Glu Thr Ser Val Met Leu Glu Leu Lys Arg Lys Met Met Tyr

106

50                   55                   60

5   Met Arg Ile Phe Ala Glu Leu Glu Gly Val Gln Glu Pro Met Ser Ile  
     65                      70                      75                      80

10                    Tyr Val Tyr Ile Ile Tyr Thr Arg Asn His Glu Val Ile Met Arg Glu  
                      85                      90                      95

15                    Phe Asn Ser Tyr Lys Arg Arg Arg His Arg Gln Asp Ala Leu Ala Tyr  
                      100                      105                      110

20                    Ala Gln Lys Ile Phe Lys Ile Leu Ser Glu His Pro Thr Phe Thr Ser  
                      115                      120                      125

25                    Gln Tyr Gln Tyr Asp Pro Val Ala Ala Leu Phe Tyr Val Lys Tyr Val  
                      130                      135                      140

30                    Tyr Glu Ser Phe Asp Ile Tyr Arg Ile Phe Ala Glu Leu Asn Val Arg  
                      145                      150                      155                      160

35                    Arg Ser Trp Glu Glu Leu Lys Ala Arg Ala Lys Lys Asp Glu Leu Ala  
                      165                      170                      175

40                    Arg Val Tyr Glu Ile Lys Cys Arg Cys Pro Ser Gln Glu Pro Met Ser  
                      180                      185                      190

45                    Ile Tyr Val Tyr Phe Glu Gln Pro Thr Glu Thr Pro Pro Gln Ile Lys  
                      195                      200                      205

50                    Val Arg Val Asp Met Val Tyr Glu Gln His Lys Ile Thr Ser Tyr Gln  
                      210                      215                      220

55                    Ala Ile Arg Glu Thr Val Glu Leu Lys Ala Arg Asp His Leu Ala Val  
                      225                      230                      235                      240

60                    Leu

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/AU02/00829

**A. CLASSIFICATION OF SUBJECT MATTER**Int. Cl. <sup>7</sup>: C07K 14/045; A61K 39/12, 39/295; A61P 31/12; C12N 15/00

According to International Patent Classification (IPC) or to both national classification and IPC

**B. FIELDS SEARCHED**

Minimum documentation searched (classification system followed by classification symbols)

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)  
 STN, Medline, CA, WPIDS, BIOSIS (keywords: polyepitop?, epitop?, cytomegalovirus, CMV, HCMV, cytotoxic T lymphocytes, CTL.)

**C. DOCUMENTS CONSIDERED TO BE RELEVANT**

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	Journal of Clinical Microbiology, Vol. 32, No. 2, Feb. 1994; Ripalti, A. et al, "Construction of Polyepitopes Fusion Antigens of Human Cytomegalovirus of ppUL32: Reactivity with Human Antibodies", pp. 358-363  See abstract and "materials and methods"	107-112, 142, 147
A	Microbiologica, Vol. 18, 1995; Ripalti, A. et al, "Construction of a polyepitope fusion antigen of Human Cytomegalovirus ppUL32 and detection of specific antibodies by ELISA", pp. 1-12.  See summary and "materials and methods"	107-112, 142, 147

 Further documents are listed in the continuation of Box C See patent family annex

* Special categories of cited documents:	
"A" document defining the general state of the art which is not considered to be of particular relevance	"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"E" earlier application or patent but published on or after the international filing date	"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"O" document referring to an oral disclosure, use, exhibition or other means	"&" document member of the same patent family
"P" document published prior to the international filing date but later than the priority date claimed	

Date of the actual completion of the international search  
17 October 2002

Date of mailing of the international search report

25 OCT 2002

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## INTERNATIONAL SEARCH REPORT

International application No.

PCT/AU02/00829

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	The Journal of Infectious Diseases, Vol. 181, 2000; Gyulai, Z. et al, "Cytotoxic T Lymphocyte (CTL) Responses to Human Cytomegalovirus pp65, IE1-Exon4, gB, pp150, and pp28 in Healthy Individuals: Reevaluation of Prevalence of IE1-Specific CTLs", pp. 1537-46  See abstract	107-112, 142, 147
A	Journal of Clinical Microbiology, Vol. 37, No. 1, Jan. 1999; Greijer, A. et al, "Molecular Fine-Specificity Analysis of Antibody Responses to Human Cytomegalovirus and Design of Novel Synthetic-Peptide-Based Serodiagnostic Assays", pp.179-188  See abstract and Table 1.	107-112, 142, 147
A	WO 94/23744 A (THE WISTAR INSTITUTE OF ANATOMY AND BIOLOGY and THE GOVERNMENT OF THE UNITED STATES OF AMERICA on behalf of THE DEPARTMENT OF HEALTH AND HUMAN SERVICES) 27 October 1994  See abstract	107-112, 142, 147
A	WO 98/21233 A (CITY OF HOPE) 22 May 1998  See abstract	107-112, 142, 147
A	WO 01/72782 A (CITY OF HOPE) 4 October 2001  See abstract	107-112, 142, 147

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/AU02/00829

**Box I Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)**

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1.  Claims Nos :  
because they relate to subject matter not required to be searched by this Authority, namely:
  
  
  
2.  Claims Nos : **1 to 106, 113 to 141, 143 to 146, 148 to 270**  
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:  
So many peptide sequences are claimed - at least 318 - that a search that is more than 3 to 4 per cent of this total is economically prohibitive.
  
  
  
3.  Claims Nos :  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a)

**Box II Observations where unity of invention is lacking (Continuation of item 3 of first sheet)**

This International Searching Authority found multiple inventions in this international application, as follows:

See attached sheets (as per attachment to "INVITATION TO PAY ADD. FEES").

1.  As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims
2.  As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3.  As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
  
  
  
4.  No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention mentioned in the claims as agreed; it is covered by claims Nos.:

107-112, 142, 147

**Remark on Protest**

- The additional search fees were accompanied by the applicant's protest.
- No protest accompanied the payment of additional search fees.

**INTERNATIONAL SEARCH REPORT**  
Information on patent family members

International application No.  
**PCT/AU02/00829**

This Annex lists the known "A" publication level patent family members relating to the patent documents cited in the above-mentioned international search report. The Australian Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

Patent Document Cited in Search Report				Patent Family Member			
WO	94/23744	CA	2160583	US	5552143	AT	140154
		CA	2012895	DE	69627695	DK	389286
		EP	389286	ES	2091221	GR	3020765
		US	5591439	WO	94/17810		
WO	98/21233	US	6074645	US	6156317	US	6251399
		AU	74814/98	CA	2308118	EP	1023319
		JP	2001519181	WO	99/19349	WO	01/72782
		AU	49243/01	EP	946592		
WO	01/72782	AU	49243/01	US	6251399	EP	946592
		WO	98/21233				

END OF ANNEX